

locus	log2ratio	lowerCL	upperCI	quantified_peptides
sp_P67762_YHBP_ECOLI	4.3	3.6	4.9	3
sp_P0A6X7_IHFA_ECOLI	3.6	2.8	7	2
sp_P21499_RNR_ECOLI	3.3	2.9	3.8	15
sp_P69913_CSRA_ECOLI	3.1	0.7	4.2	2
sp_P0A7L0_RL1_ECOLI	3	2.6	3.5	19
sp_P0A8R0_RRAA_ECOLI	2.8	1.7	7	2
sp_P0AEJ2_ENTC_ECOLI	2.8	2.2	3.6	5
sp_P0A6X3_HFQ_ECOLI	2.3	1.8	2.8	5
sp_P0A7V8_RS4_ECOLI	2.1	1.7	2.5	9
sp_P0AG67_RS1_ECOLI	2.1	1.8	2.4	30
sp_P37765_RLUB_ECOLI	2.1	1.4	7	3

Numerator: Scrambled (Heavy)
Denominator: S12 (light)

description

UPF0306 protein yhbP OS=Escherichia coli (strain K12) GN=yhbP PE=3 SV=1 [MASS=16776]
Integration host factor subunit alpha OS=Escherichia coli (strain K12) GN=ihfA PE=1 SV=1 [MASS=11354]
Ribonuclease R OS=Escherichia coli (strain K12) GN=rnr PE=1 SV=2 [MASS=92109]
Carbon storage regulator OS=Escherichia coli (strain K12) GN=csrA PE=1 SV=1 [MASS=6856]
50S ribosomal protein L1 OS=Escherichia coli (strain K12) GN=rplA PE=1 SV=2 [MASS=24730]
Regulator of ribonuclease activity A OS=Escherichia coli (strain K12) GN=rraA PE=1 SV=1 [MASS=17360]
Isochorismate synthase entC OS=Escherichia coli (strain K12) GN=entC PE=1 SV=1 [MASS=42932]
Protein hfq OS=Escherichia coli (strain K12) GN=hfq PE=1 SV=2 [MASS=11166]
30S ribosomal protein S4 OS=Escherichia coli (strain K12) GN=rpsD PE=1 SV=2 [MASS=23469]
30S ribosomal protein S1 OS=Escherichia coli (strain K12) GN=rpsA PE=1 SV=1 [MASS=61158]
Ribosomal large subunit pseudouridine synthase B OS=Escherichia coli (strain K12) GN=rluB PE=1 SV=2

[MASS=32711]